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### FIG. 2A.

	<b>-13</b>								
	M	D	W	T	W	R	F	L	
TCTAAAGAAGCCCCTGGGAGCACAGCTCATCAC	ATG	<b>GAC</b>	TGG	ACC	TGG	<u>AGG</u>	TTC	CTC	58
	STG	NAT.	<u> </u>	_					

F V V A A A T G V Q S Q M Q V V Q S G A

TTTGTGGTGCAGCTACAGGTGTCCAGTCCCAGATGCAGGTGCAGTCTGGGGCT 118
FR1⇒

10 15 20 25 E V K K P G S S V T V S C K A S G G T F GAAGTAAAGAAGCCTGGGTCCTCGGTGACGGTCTCCTGCAAGGCATCTGGAGGCACCTTC 178

30 31 32 33 34 35 40 45
S N Y A I S W V R Q A P G Q G L E W M G
AGCAACTATGCTATCAGCTGGGTGCGACAGGCCCCTGGACAAGGGCTTGAGTGGATGGGA 238
CDR1⇒ FR2⇒

50 51 52 a 53 54 55 56 57 58 59 60 61 62 63 64 65
G I I P L F G T P T Y S Q N F Q G R V T

GGGATCATCCCTCTTTTTGGTACACCAACCTACTCACAGAACTTCCAGGGCAGAGTCACG 298
CDR2⇒
FR3⇒

70 75 80 82 a b c 83 85 I T A D K S T S T A H M E L T S L R S E ATTACCGCGGACAAATCCACCAGCACAGCCCACATGGAGCTGACTAGCCTGAGATCTGAG 358

f g h i k 101 102 105 110 115 R V G W F D P W G Q G T L V T V S S A S CGGGTTGGCTGGTCCCTCGGGCCAGGGCACCCTGGTCACCGTCTCCTCAGCCTCC 478  $C_{\mu}1\Rightarrow$ 

120 125 130 133 135
T K G P S V F P L A P S S K S T S G G T
ACCAAGGGCCCATCGGTCTTCCCCCTGGCACCCTCCTCCAAGAGCACCTCTGGGGGCACA 538

140 145 150 154 156 162 A A L G C L V K D Y F P E P V T V S W N GCGGCCTGGGCTGCCTGGTCAAGGACTACTTCCCCGAACCGGTGACGGTGTCGTGGAAC 598

165 169 171 175 180 182 S G A L T S G V H T F P A V L Q S S G L TCAGGCGCCCTGACCAGCGGCGTGCACACCTTCCCGGCTGTCCTACAGTCCTCAGGACTC 658

185 190 195 200 203 205 Y S L S S V V T V P S S S L G T Q T Y I TACTCCCTCAGCAGCGTGGCCCTCCAGCAGCTTGGGCACCCAGACCTACATC 718

## FIG. 2B.

C TG		N AC	21 V GT	N	-	H CAC	K CAA		)	215 S AGC	N	T CAC	K CAA	_	Ţ	220 D GAC	K	222 K GAA	V AGT	E	GCC	-	K	232 S TCT	778
C TG	I TG/	D	23! K AA	Т		H CAC	T AC		•	240 P CCA	P CC(	C GTG(	P CCC	A AGC C <sub>H</sub>	A		E	L ACT(	L CCT	G GGG		}	P CG1	S FCA	838
V GT	F CTT	?	255 L CTC	F	CC	P CC	P CC.	K AAA		60 P CC	K AAC	D GA(	T CAC	L CCT	,	265 M ATG	I AT(	S CTCC	R CCG	T GAC	27 F CCC	•	E AG(	V STC	898
T AC	C ATG		275 V GTG	V		V TG	D GA(	V CGT	•		H CAC		D AGA(		1	285 E GAG		K CAAC		N CAA	29 W	1	Y	292 V STG	958
299 D GA	G	ř	•					N ГАА		Α		T				R		E	Q CAC	Y			S		1018
Y TAC	32 R CCG	2	V GTG	_	CA	_ <del>_</del> _		32 L CCT		_	V FTC	L CTG	H CAC	•	-	D BACT	W TGG	L CTG	N AA7	33; G GG(	K	_		Y 'AC	1078
K AAC	34 C TG		K AG	V GT		s ÇC	N AA(	34 K CAA		A CCC	L CTC	P CCA	A .GC(			I	E GAG	K AAA	T	355 I CAT(	S	]		A CC	1138
360 K AAA	G	GC	63 Q CAG	P				P ACC		_	V	370 Y TAC	T				P	375 S TCC	R	D	378 E [GA	]	L	T .CC	1198
K AAC	N AA		85 Q AG	V GT(		S GC(		T FAC		_	L TG		K AAA	G \GG(			Y CAT	P	S AGC	D GAC	40 I CAT	Ā	4	02 V TG	1258
405 E GAG	W	GG	E	08 S AG(	]	10 N AT(	G	14 Q GCA	•	P	E AG.	N AAC	N AAC	Y TAC		20 K AGA	T	T ACG	P CCI	P	42 V CGT	I	ւ rgg	D AC	1318
S	D		G	S		F	F		•	Y	S		L				D	445 K AAG	S				Q .		1378
																		465 N AAC							1438
	_		_			_		G GG'				GTG	CGA	CG	GC	CGG	CA	AGC	CCC	CGC	СТС	cco	CGG	GС	1498
TCT	,CG	CG	GT	CGC	CAC	CGA	4GG	AT(	GC'	ГТG	GC	ACG	TAC	CCC	CG	TGI	CAC	ATA	CTT	CC(	CGG	GC	GCC	CA	1558
GCA	TG	GA	AA'	ΓΑΑ	AA(	GC/	ACC	CA	GC	GCT	GC	сст	GGG	CC	CC	TGC	CGA	AAA.	AAA	AAA	AAA.	AA <i>A</i>	AAA	A 1	617

## FIG. 3A.

	CAA	GAC	GCA	.GCG	CTC	TC	GGG!	ACGT	'CTC	CAC	-1 M CAT	A	W CTG		L TCT	, L	i I	L CCT	T	5.8
												GNA		<u></u>	<u> </u>	GOZ	<u> </u>	<u> </u>	CACC	. 30
L CT	L CCT	T CAC	TCA		T CAC	AGG		· · · · · · · · · · · · · · · · · · ·		_	_			5 T GAC	v	_		9 S CTC	11 V CGTG	112
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S	G	S	15 P	G	ଚ	S	. T	20 T	т	S	•			26	-	d	e	f	28	
TC	_	GTC	_				_	_	_		_			T AAC	N CAA	N CAA	TGA	V TGT:	G TGGG	178
	<b></b>											CD	R1⇒							
29 S	30 Y	31 N	32 L	33 V	34 S	35 W		Q	Q	Н	40 P	G	K	Α	P	45 K	I	М	I	
<u>AGI</u>	'TA'	ΓΑΑ	CCT	<u>rgt(</u>	CTC	CTG FR		CCA	GCA	GCA(	CCC	AGG	CAAA	AGC	CCC	CAA	AAT	CATO	GATT	238
	50	51	52	53	54	55	56				60					65				
Y rat	E	V	S	K	R	P	S	G AGG(	V تون	S	N	R	F	S	G	S	K	S	G CGGC	0.00
• • • •	CDI	22⇒	<u>UAU 1</u>	AAC	rcac	<u>rcc</u>		FR		110	LAA	CGC	J I'I'C	TO	ruu	Free	JAA	JTC1	rGGC	298
	70					75					80					85				
N AAC	T AC	A GCC	S CTCC	L CTG	T ACA	I ATO	S CTC:	G TGG0	L CTO	Q CCAC	A GGCI	E GAG	D GAC	E GAC	A GCT	D TGAT	Y TAT	Y TAC	C ·	358
39 C	90 S	91 V	92 A	93 G	94 S	95 Y	a T	96 V	97 V	F	G	100 G	•	æ	17	7	m	106	a	
GC DR	TCA	TAT		_	_	_	•	_	-	TTT	cocc		G .GGG	ACC	AAA	CTC	ACC	v CGTC	CTA	418
	J₹									FR4	ŀ⇒									
07 G	Q	P	110 K	A	Α	P	S	115 V	T	L	F	P	120 P	S	S	E	E	125 L	ရ	•
GT(	CAG C <sub>1</sub> ⇒	CCC	AAG	GCT	GCC	CCC	CTCC	GTC	ACT	CTG	TTC	CCG	CCC	TCC	TCT	'GAG	GAG	CTT	CAA	478
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A	N	K	Α	T	L	V	C	L	I	S	D	F	140 Y	P	G	A	v	145 T	v	
																			GTG	538
A	W	K	150 A	D	s	s	P	155 V	K	Α	G	v	160 E	Т	Т	Т	p	165 S	К	
CCI	rgg.	AAG	GCA	GAT	AGC.	AGC	CCC	GTC	AAG	GCG	GGA	GTG	GAG	ACC	ACC	ACA	CCC	TCC	AAA	598
ျ ( ဂ	88 S	170 N	N	ĸ	v	Δ	175	c	e	v	¥	180	•	<b></b>	_	_	185		•	
AAA	GC.	AAC	AAC	AAG	rac	GCG	GCC	AGC.	AGC	TAT	CTG.	AGC	CTG.	ACG	CCT	GAG	CAG	W TGG.	K AAG	658
~		190	S	_			195				2	00	203		205					
S CCC	H AC	R AGA	S AGC	Υ CAC	S AGC'	C TGC	Q CAG	V GTC	T ACG	H Cat	E GAA	G GGG.	S AGC	T ACC	V GTG	E GAG	K AAG	TACA	V GTG	718

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## FIG. 3B.

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	CDR1  ********  27 a b c  3TCAG A A	 GAGCCTTTAT CDR2	TGCATCCAGTTTG	G-TTACCG- G-TTACCG- GACCGA
	GGAGACA GT	GGC	CAGAAAC	CC-TAG  CTAG  CG
FIG. 4A.	CAGATGACCCAGTCTCCATCCTCCTGTC	TGT	AGCATTAGCAATTAAATTGGT GGGG	GAGTC GCGGCC GTAGC-CGCC TCAAGCAAACA-AAGGTGC-CGCC
	この45のエエエ・	CynoK20 Rabbit Mouse	- A CH 10 Ch .	Cynokli Cynokli Cynokli Cynokli Rabbit Mouse

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FIG. 5A.

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	, <u>.</u> <u>.</u> <u>.</u> <u>.</u> <u>.</u> .
Human Cynomolgus Rabbit Mouse	110 ACTGTGCACCATCTTCATCTTCCCGCCATCTGATGAGCAGTTGAAATCTGGAACTGCCTCTGTTGTGTGCCTGCTGAATAAC GTT
Human Cynomolgus Rabbit Mouse	140 TTCTATCCCAGAGGCCAAAGTACAGTGGAAGGTGGATAACGCCCTCCAATCGGGTAACTCCCAGGAGTGTCACAGAGCAGGACAGC
Human Cynomolgus Rabbit Mouse	170 AAGGACAGCACCTCAGCAGCACCCTGACGCTGAGCAGACTACGAGAAACACACAAAGTCTACGCCTGCGAAGTCACCCATAAATT-ACCAC-GCGCTAGGGT G-AT-T-TA
Human Cynomolgus Rabbit Mouse	200 CAGGGCCTGAGCTCACAAAGAGCTTCAACAGGGGAGAGTGTTAGAG-GA-GTCCTTC AACATCA-CTA-TGTCAATAAT

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F G. 7,   120					•	
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FIG. 7.  110  120  130  140  140  140  140  140  140  14						
F G, 7,   110						
110  110  110  120  130  140  140  140  140  140  140  14						
110  TVAAPSVFIFPPSDEQLKSGTASVVCLLNNF  120  TVAAPSVFIFPPSDEQLKSGTASVVCLLNNF  140  LOSGNSDEQLKSGTASVVCLLNNF  160  LOSGNSDEQLENTTEFFT  170  LOSGNSDEXT  180  LOSGN		0				
110  T V A A P S V F I F P P S D E Q L K S G T A S V V C L L N N  130  T V A A P S V F I F P P S D E Q L K S G T A S V V C L L N N  140  D P I - T - L		<b>←</b>		•		
110  T V A A P S V F I F P P S D E Q L K S G T A S V V C L L N  110  T V A A P S V F I F P P S D E Q L K S G T A S V V C L L N  D P I - T - L L						
110  120  130  1 VAAPS VFIFP SDEQLKSGTAS VCLL  Slgus A						
110  120  130  140  150  160  160  160  160  160  160  16						
110  120  17			0			
110  T V A A P S V F I F P P S D E Q L K S G T A S V V V A A P S V F I F P P S D E Q L K S G T A S V V V D P I I I L L L L L L L L L L L L L L L L			- ←			
110  1 V A A P S V F I F P P S D E Q L K S G T A S V O D D D D L V O D D D D D D D D D D D D D D D D D D						
110  T V A A P S V F I F P P S D E Q L K S G T A S S T Y T L L L L L L L L L L L L L L L L L						
110  120  120  130  140  150  160  160  160  160  160  160  170  17						
110  T V A A P S V F I F P P S D E Q L K S G T O 1 D P I - T T - L L - T - T - E D - V - T - T - E D - V - T - T - E D - V - T - T - E D - V - T - T - E D - V - T - T - E D - V - T - T - E - T - T - T - E - T - T - T						
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110  120  110  1 V A A P S V F I F P P S D E Q L K  1 V A A P S V F I F P P S D E Q L K  2 D P I - T - L L B D T  A D T - S T  A D T - S T  1 Q S G N S Q E S V T Q Q D S R D S  2 1 Q S G I E N - T - P - S P E - C  E R Q N G V L N - W - D T  2 2 0 Q L S S P V T K S F N R G E C  2 1 Q G L S S P V T K S F N R G E C  2 1 D						
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FIG. 7.  110  T V A A P S V F I F P P S D E O  120  T V A A P S V F I F P P S D E O  120  D P I T - L L B D  A D T - E L B D  A D T - S B D  160  L O S G I E N - T - P C S P E  E R O N G V L N - W - D  200  O G L S S P V T K S F N R G E C  210  O G L S S P V T K S F N R G E C  210  O G L S S P V T K S F N R G E C  210  O G L S S P V T K S F N R G E C  210  O G L S S P V T K S F N R G E C  210  O G L S S P V T K S F N R G E C  210  O G L S S P V T K S F N R G E C  210  O G L S S P V T K S F N R G E C  1			0			
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110  T V A A P S V F I F P P S D  110  T V A A P S V F I F P P S D  120  D P I - T - L L A  A D - T - T - L L B  A D - T - T - L L B  160  L Q S G N S Q E S V T Q Q D  190  L Q S G N S Q E S V T Q Q D  200  S P V T N S P P S N R G  210  Q G L S S P V T K S F N R G  210  Q G L S S P V T R S F N R G  210  Q G L S S P V T R S F N R G  210  Q G L S S P V T R S F N R G  210  Q G L S S P V T R S F N R G  210  Q G L S S P V T R S F N R G  210  Q G L S S P V T R S F N R G  210  Q G L S S P V T R S F N R G  210  Q G L S S P V T R S F N R G  210  Q G L S S P V T R S F N R G  210  Q D L S S P V T R S F N R G  210  Q D L S S P V T R S F N R G  210  Q D L S P V T R S F N R G  210  Q D L S P V T R S F N R G  210  Q D L S P V T R S F N R G  210  Q D L S P V T R S F N R G  210  Q D L S P V T R S F N R G  210  Q D L S P V T R S F N R G  210  Q D L S P V T R S F N R G  210  Q D L S P V T R S F N R G  210  Q D L S P V T R S F N R G  210  Q D L S P V T R S F N R G  210  Q D L S P						
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FIG. 7.  110  T. V. A. A. P. S.  S. D. G. M. S.  D. J. J.  E. M. S. G. M. S.  C. J. S. S. P.		E I H U	0			
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